## P03001-seq list.ST25

## SEQUENCE LISTING

<110> Chen, Chao-Ying

<120> A NOVEL NUCLEIC ACID ENCODING BETA-1,3-GLUCANASE FROM LILY

<130> p03001

<160> 2

<170> PatentIn version 3.2

<210> 1

<211> 337

<212> PRT

<213> lily

<400> 1

Met Ala Ala Gln His Ile Ile Ser Met Ala Ala Met Ala Ser Leu Leu 1 5 10 15

Val Val Leu Ser Ala Ile Pro Arg Gly Val Glu Ser Ile Gly Val Cys
20 25 30

Asn Gly Met Asp Gly Asp Asn Leu Pro Gln Pro Ala Asp Val Val Asn 35 40 45

Leu Tyr Lys Ser Asn Asn Ile Ala Gly Met Arg Leu Tyr Ser Pro Asp 50 60

Gln Ala Thr Leu Gln Ala Leu Gln Gly Ser Asn Ile Tyr Leu Ile Leu 65 70 75 80

Asp Val Pro Asn Ser Asp Leu Gln Asn Ile Ala Ser Asp Gln Ser Ala 85 90 95

Ala Thr Asn Trp Val Gln Thr Asn Val Gln Ala Tyr Pro Asn Val Ala 100 105 110

Phe Arg Tyr Ile Ala Val Gly Asn Glu Val Ile Pro Gly Gly Gln Ala 115 120 125

Gln Tyr Val Leu Pro Ala Met Asn Asn Ile Gln Ser Ala Leu Ser Ser 130 135 140

Ala Gly Leu Gln Asn Ile Lys Val Ser Thr Ser Val Ser Phe Gly Val 145 150 155 160

Val Gly Thr Ser Tyr Pro Pro Ser Ala Gly Ser Phe Ser Ser Asp Ala 165 170 175

## PO3001-seq list.ST25

P03001-Seq 1131.3123	
Ser Ser Thr Leu Gly Pro Ile Ile Gln Phe Leu Ala Ser Asn Gly Ser 180 185 190	
Pro Leu Leu Ala Asn Ile Tyr Pro Tyr Leu Ser Tyr Ala Gly Asn Ser 195 200 205	
Gly Ser Ile Asp Leu Ser Tyr Ala Leu Phe Thr Ala Ser Gly Thr Val 210 215 220	
Val Gln Asp Gly Ser Tyr Ala Tyr Asn Asn Leu Phe Asp Ala Met Val 225 230 235 240	
Asp Ala Leu Tyr Ser Ala Leu Glu Ser Ala Gly Gly Pro Asn Val Pro 245 250 255	
val val val Ser Glu Ser Gly Trp Pro Ser Ala Gly Gly Thr Ala Ala 260 265 270	
Thr Val Ser Asn Ala Gln Thr Tyr Asn Ser Asn Leu Ile Asn His Val 275 280 285	
Gly Gln Gly Thr Pro Lys Arg Pro Gly Ala Ile Glu Thr Tyr Ile Phe 290 295 300	
Ala Met Phe Asn Glu Asp Gln Lys Gln Pro Gln Gly Ile Glu Asn Asn 305 310 315 320	
Phe Gly Leu Phe Tyr Pro Asn Glu Gln Pro Val Tyr Ser Ile Ser Phe 325 330 335	
Thr	
<210> 2	
<211> 1011	
<212> DNA	
<213> lily	
<400> 2	
atggcagctc agcacatcat ctccatggct gccatggcat ccctccttgt agtactctcg	60
gcaatcccga gaggcgtgga atccattggg gtctgcaatg gaatggacgg tgacaacctc	120
ccccagcccg ccgacgtcgt caacctctac aagtccaaca acatagctgg catgcgactc	180
tacagccccg accaagccac tctccaggcc ctccagggct ctaacatcta cctcatcctc	240
gacgtcccca actccgacct ccaaaacatt gcctccgacc aatccgccgc caccaactgg	300
gtccaaacca acgtccaagc ctacccaaac gttgccttcc gatacatcgc cgtcggaaac	360

## P03001-seq list.ST25 gaagtcatcc ccggcggcca agctcagtac gtcctccag ccatgaacaa catacagtcc 420 gccctctcct ctgccggcct tcagaacatc aaggtctcca catcagtctc cttcggcgtc 480 gtcggtacct catatccccc ctcagctggc tccttctctt ccgatgcatc gtcgacattg 540 ggtccaatca tacagtttct agccagcaat ggctccccat tacttgccaa catctacccc 600 tacttgagct atgctggcaa ctccggatcc atcgacctct catacgccct ctttactgca 660 tctggtacag tcgtacagga cgggtcctac gcttacaaca acctcttcga tgccatggtc 720 780 gagagtggct ggccgtcagc gggcgggaca gcggcgacgg tgtctaatgc gcagacttac 840 aattccaatt tgatcaacca tgtgggtcag gggacgccga agaggccagg ggcgattgag 900 acctacatat ttgccatgtt caacgaggat cagaagcagc cgcaagggat tgagaataac 960 tttgggctgt tttaccctaa cgaacagcct gtctattcga tcagcttcac t 1011